

FIG. 1A

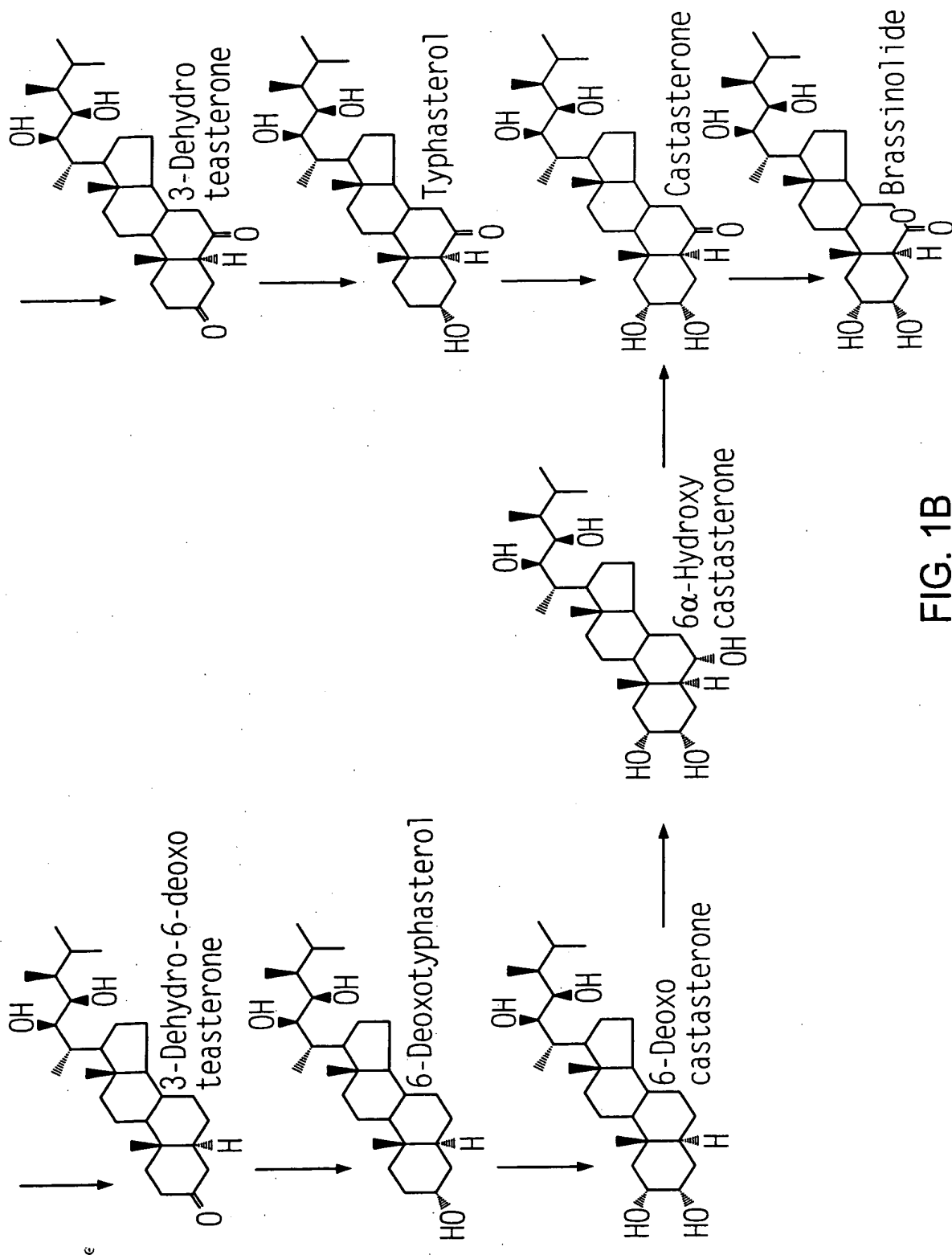


FIG. 1B

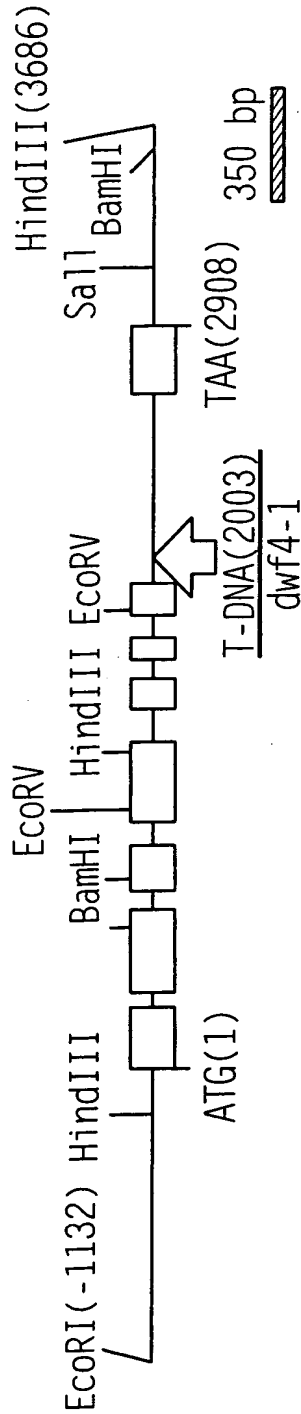


FIG. 2A

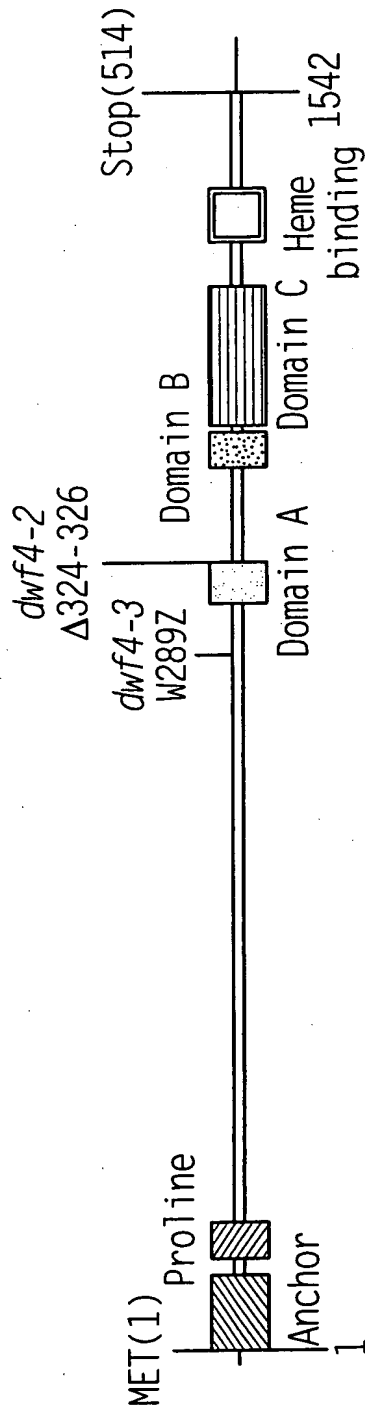


FIG. 2B



1 20 30 40 50 60 70
MFEITEHHTLLPLLL
MAFF7
MAFF
MAFF
MLGVGMAAVALLGAVALLADAAARRAHWWYREAALVGAVALVVDAAARRAHGWYREAALGAARRA
MGLYTLMTFLCTIVLPVLFLAVALKLEWMLMIRVDPNCRS
MALPDLAMETWLLAVSLVLLYLYGTHSHGLFKK
- - - - -

DWF4<CYP90B>
CPD<CYP90A>
Tomato<CYP85>
Cyanobacteria<CYP120>
Maize<CYP88>
Zebrafish<CYP28>
Human<CYP3A3>
CONSENSUS

ANCHOR REGION

80 90 100 110 120 130 140 150
NLPPGKSGWPFLLGETIGYLKPPYTATTLGDFMQQHVSXKYG
GLPPGSLGLPLIGETIFQIGAYKTENPEPFI DERVARYG
NLPPGTMGWPLFGETTEFLKL - - - - -GPSFMKNQARARYG
PLPPGDFGLPWGETLNF - - - - -LNDGDGKKRQOQFG
RLPPGEMGWPLVGMWAFRAFKSGKPDFAIAFVRRFGRTGVYRSFMFSSTPLVTTAEGCKQVLMDD - - - - -AFVTGWPKATVAL
PLPPGTMGWPLFGETLQLI - - - - -LQRRKFLRMKQKYG - - - - -CIYKTHLFGNPTVRVMGADNVRQILLGEH - - - - -KLVSVQWPAASVRTI
LGIPTPTPLPFLGNI LSYHKGFCM - - - - -FDMCHKKYGKVGWGYDGO - - - - -OPVLAITDPDMIKLVLVKECYSVFTNREPFGPVGF
- - - - -

PROLINE

160 170 180 190 200 210 220 230 240
LGKWSMLVLVVGDMRRDRMRSLSLNFI LSHARLRTILKKIDVERIRHTLFFVLDVSW
LGKHSLL LMKGSLMKRMHSLTMSFANSSI KDHLMLDI DRILVRFRSHLTDVSW
LGKCNIAAVNGSAMKYMGRGALLSLISPTMIRDQLLPKIDEFMRSHLTNW - - - - -SSR-VLL - - - - -MEEAKKITFELTVKQLMSFDPGEE
LGNALATQMGEIMRSRRIKLYQAFLP-RTLDSYLPKMDGIVQGYLEOW - - - - -GKANEVIWYPOLRRMTFDVAATLFGM - - - - -EST-SL
VGPRSFVAMPYDEMRRRIKLTAAPIINGFDALTGYLPFIORTVTSSLRRAW - - - - -ADHGGSVFELTELRRMTFKIIVQIFLG - - - - -GADQAT
LGSDTLSNVHGVQHKKKKAI MRAFSR-DALEHYIPVIOQEVKSALQEW - - - - -LQKDSCLVLPVPEMKLMERIAMRILLGFEPEQIK
M-KSAISIAEDEEWKRLRSLSPFTTSGLKE-MVPI LAQYGDVLVRNLRRERETGKPVTLKDVFGAYSMDVITSSFGVNVDSL
- - - - -

250 260 270 280 290 300 310 320
TE - - - - -QLKKEYVTFMKGVVSAPLNLPLGTAHKAHQSRATILKFI ERKMEERKLDI KEEDOE EEEVKT EDEAE MSKSDHVRKORITDD
SE - - - - -SLRKEYLLVIEGFFSLPLPLFSSTTYRKAIQAR - - - - -RKVAE - - - - -ALTVVMKRRREEE - - - - -E EGAEE - - - - -RK KDM
AQ - - - - -EFMSEFFNLVLTSLPLSLPI NLPTNYHRGFOQARKIIVNL LR - - - - -TLFEERRASKEI - - - - - - - - - - -QHD
SQNPQLFPWFETIYQGLFSLPIPLPNTLFGKSQARALLAELEKIKAR - - - - -QQQPPSEE - - - - - - - - - - -DA
TR - - - - -ALERSYTELNYGMRAAI NLPGFAYRGALRRRLVAVLO - - - - -GVLDERRAARAKVSGGV - - - - - - - - - - -DM
TDEQLVEAFEEMIKNLFSPLIDVPFSGLYRGLRARNFIHSKI EENIRKK - - - - -I QDDDNENE - - - - - - - - - - -QKYKDA
NPQDPLVENTKLLRFDPLPFLSI TVF - - - - -PFLPILE - - - - -VLNICVFPREVTNFLRKAVRMKESRLED TQKHRVD
- - - - -

FIG. 3A

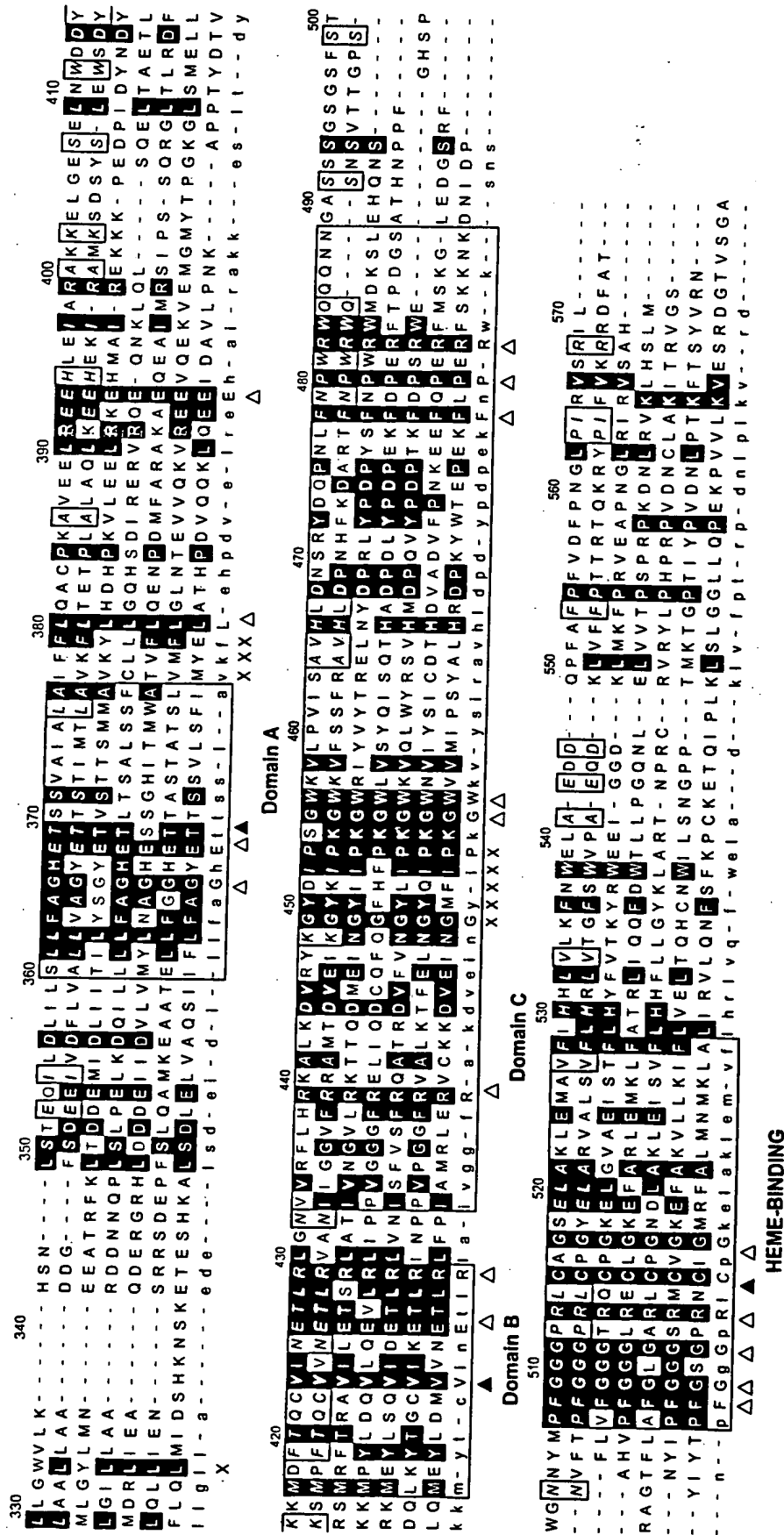
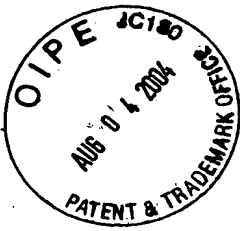


FIG. 3B

HEME-BINDING

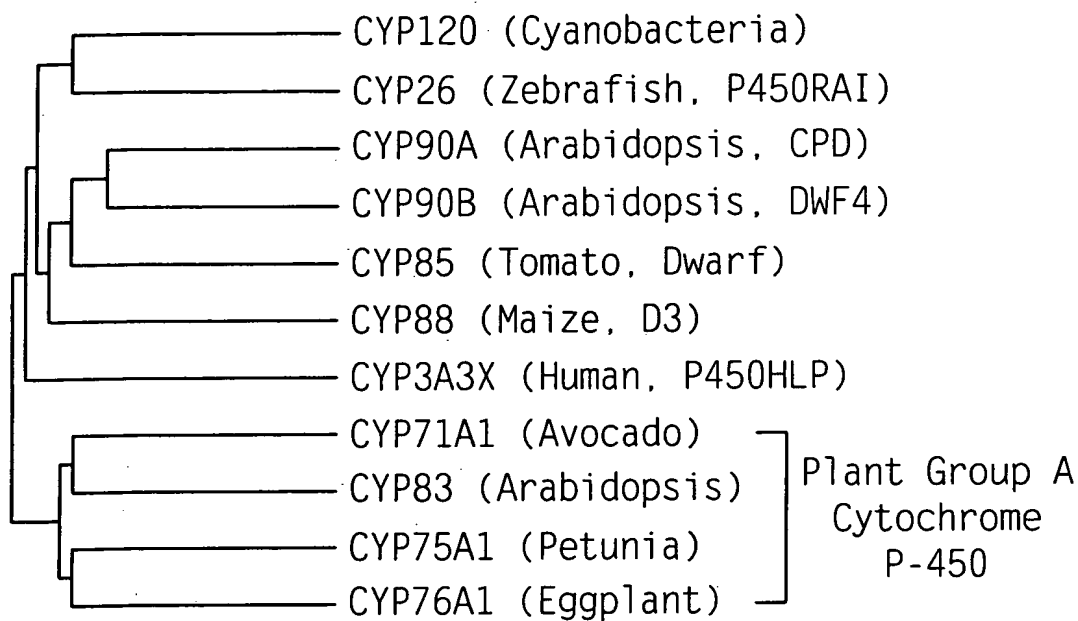
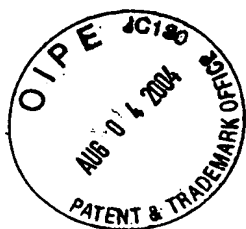


FIG. 4

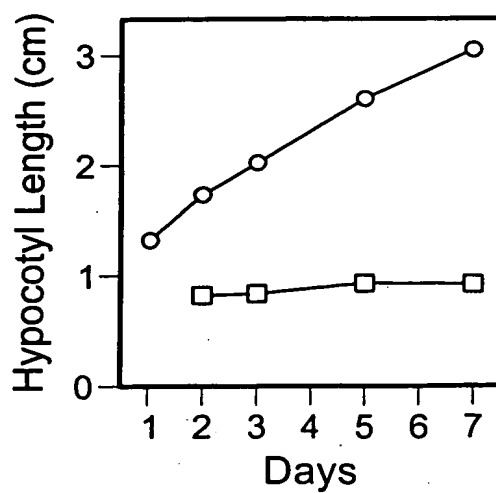


FIG. 5

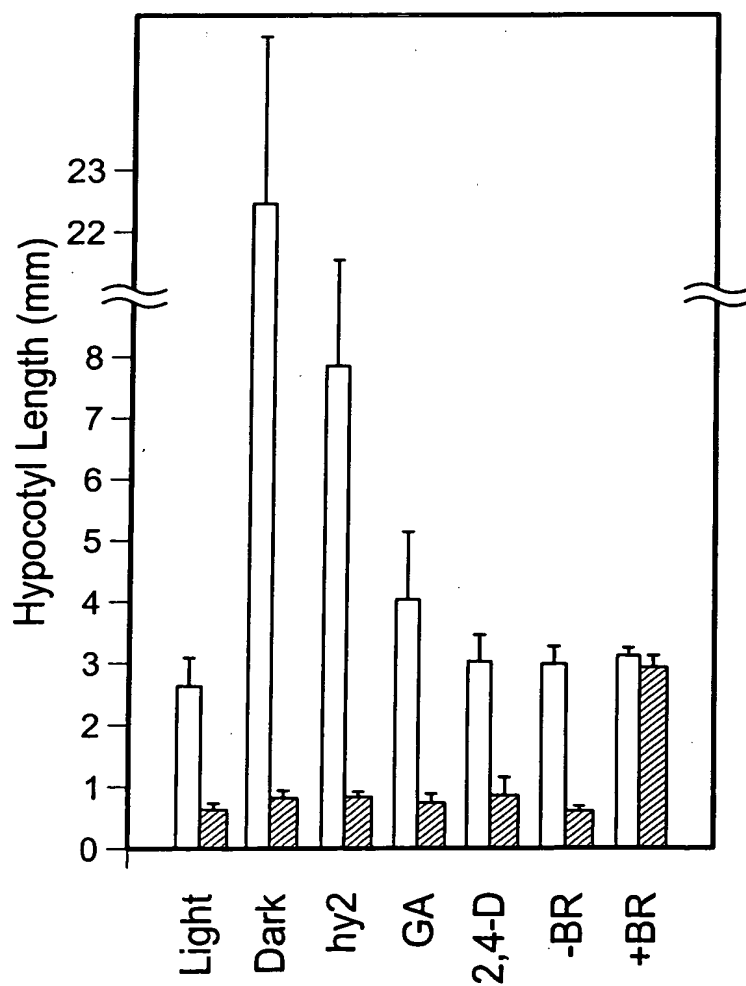
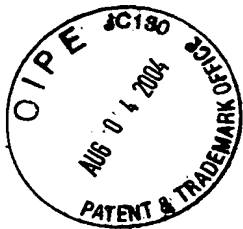


FIG. 6

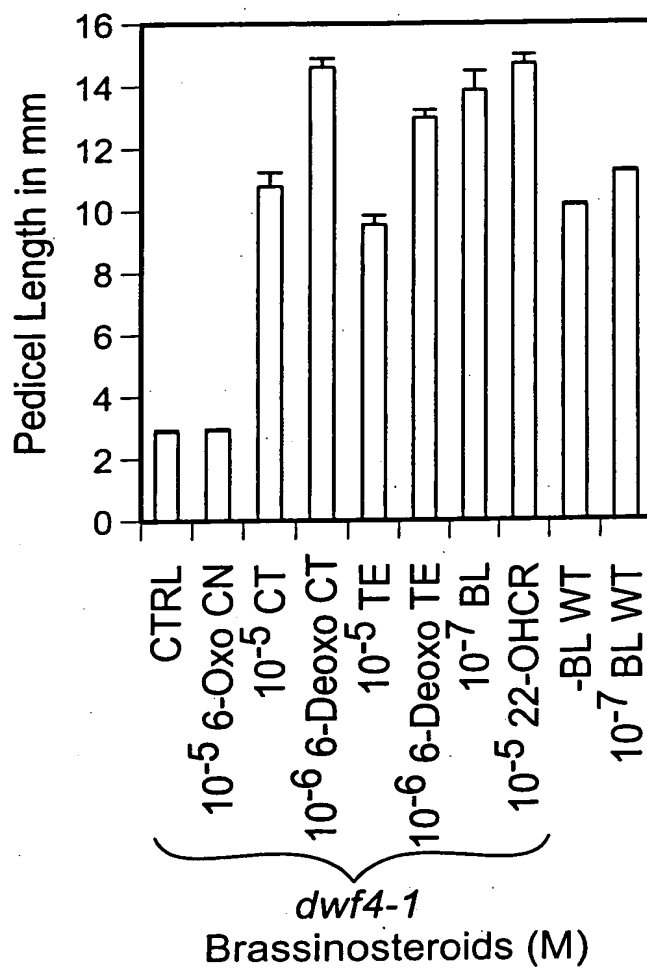
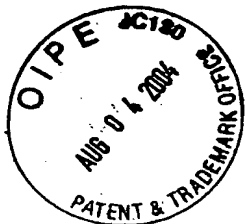
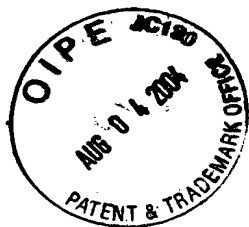


FIG. 7



Increase of Stem Growth

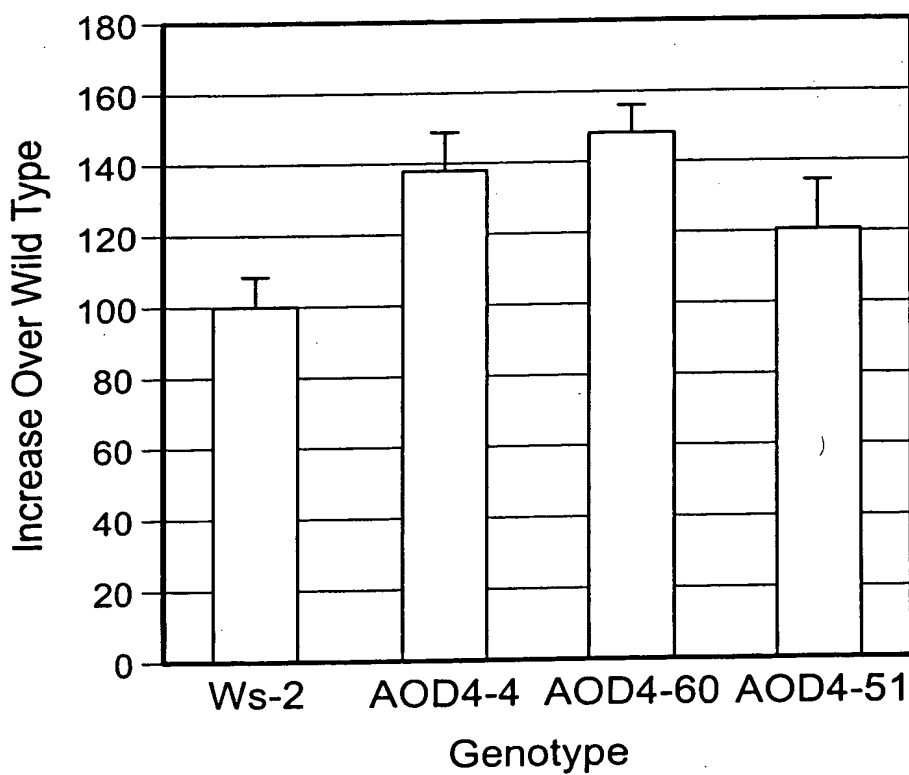
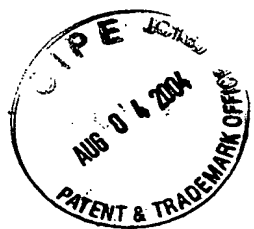


FIG. 8



Increased Seed Production Due to DWF4 Overexpression

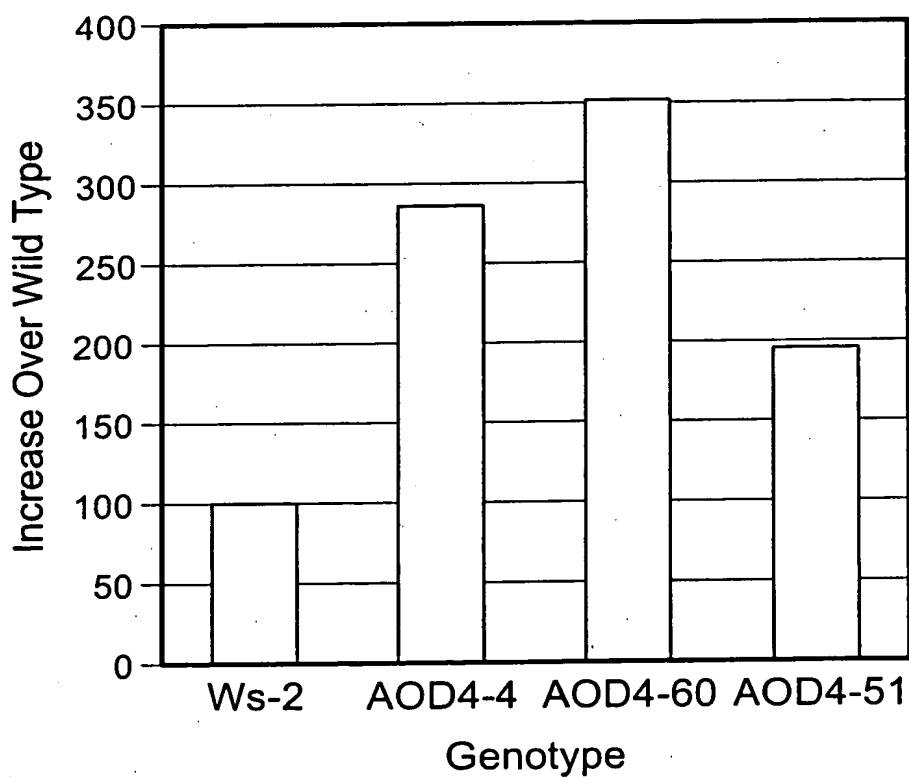


FIG. 9



1 ATGTGGTATTATTGTTGGGTTGGTTTGAGCTACAATAATAAATTCGIGTTTCIGGT 60
61 TATTCTGTTACATGATTGAGTTTGGTTCTCAATTTGGATTCCAAGATAATTAATATT 120
121 AAAATTCATTTAAATATTTACAAGTAATTAATTATCTTTACATTGTATTGTTATAACAA 180
181 AATACTAICTTTGGTATATGAGAAAATATGGAGTTTGGAAATTTATAATAATAAGGAAA 240
241 TAATCGATTCCATTGGTTGGATTACACAGTTAAGTTTTTGIGTTTCITTTGTTATAATGT 300
301 ATATGAGTAAATCAAAAAGAGTATTGATTGAAGTGTAACATAATTCGTTATGACCCCCA 360
361 AAAAAAAAAAAAAACAAACAAACCCCCCGATATAGTTTTTGGTTCTGGATT 420
421 AGGTTTATTGATCATAATTACATGCATCATTTCTTTGATTACTAIGAAGATTTTCTTAC 480
481 CAATTAAATTCGAATTCATACTCTTGATTATTAAATTAACGAGTGTGAATATCC 540

FIG. 10A



541 GTTATCGATCACTCCAATCATGATTATGATTCTTGTGCTAATCCAGCAAATTATTAAACA 600
601 AGAGTATTGAGAAAAAACCAGAAAATAAGAAAGGAAAGAGTAGTGACCCATGGAGTATG 660
661 TGAATAATTATCAAGAGAGAATAAGAGATGACAACCAAAAGGTTGTGGAATAATGGTCCCT 720
721 GCCAGCTTTCTCTCACAAATCAATAATCGACCCTATTTGGATTTTCTGGATATTCGTTAAAA 780
781 TTTCGATAACGATTGTGAAAAAATTTTATTTGTTAGCTGATCTCAATATTATGTTCCTCA 840
841 GGTATTGTCATAATCTTCGTGTTAAAGCATATTTTGTCTTTCTTTTGTTCGTTTCTCT 900
901 TAACTATATATTATCGCGGATATATGATAACAATGATATATCACAAAACAATTGTCTGGG 960
961 ACCATTTTGAATAAACTTTTTCTCAAAACATTACGGGACACTGGACTCGACCCTTAAATA 1020

FIG. 10B



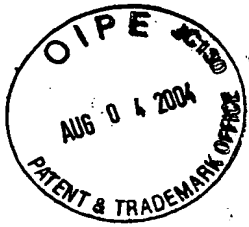
1021 CGATTTTACAGCGTCACTAGTTGAGATTACTAGCATAAAGGACCCGTTCAAGC 1080
1081 TATTTATACAAAGTTACAAACTGAATATAGCTTGAAATCCTTTAGAAAATTTTGGAAATTA 1140
1141 CCGGTTGTTATGTAATATAGATTTAGTGGTAACAAATATGTTAATCAATTAGTGGTCA 1200
1201 ACATATACATAATTCCTTACAGAAAAACAACTTAAGAGAAGTTAACATATCCATATAT 1260
1261 GGGTATGCTATACCTTTCACGTA TGCTATACTAGAGACTAAAGAATAGTTATGTGATGTC 1320
1321 GATAAATGAAATTCACACGCGTGGTAATAATTATGGGACCGTATGTTACGATCAGTCAA 1380
1381 ATATCATTCCTTGGTGGTCAACAATAAAAAACAACAAAAAAGAAAAACGATTTTTT 1440
1441 CTGGATTCCATTCAATGATCTAAAATGCATAGATCTTTTGGGTTACAGTTTCGAAGTCC 1500
1501 TCTACAAGCGGTAAACCATCTGCAACTATTAAATTGCTTTCCTTTAATGCATCTTTAACAT 1560

FIG. 10C



1561 ATTTATTGTTAGTTGGAATTTAATAAGAGCGAACTTGTAACATTACAATAATTTATATTAG 1620
1621 ATACTAGTAGTGATTATTCCAAATACATACTTTGGATGTTTAACTTAAICTTGTTTCT 1680
1681 TCCTACGGTATAAATATTAATCATCGAGGTAAAAAAGTTTTGTCCTTATTTTCGGGATGC 1740
1741 ATGAAGGATAAACCTAATGACITTTAATTTTTTGAAAAATGTAACCCCTTTTACTCATAGATT 1800
1801 AATTACCGTAIGTTTTTGTGGCCATAATGACAGCCCTCTACAACCTGTGATAGTCAATTTTT 1860
1861 TCTGCAAAATATTAAATTAGGAATTCATGCTACTATCAATAGAGAACAGCTGAGTATT 1920
1921 ACATTTTAAATTAAGACACAAAATTTTTGAAAAATGTTATAATTTCTAACAATATTATTAA 1980
1981 AATATGATGCCATATAATGTAATTCCTAATGTTCTTAAAAATATTTTTTTTATATTAGTTA 2040
2041 TAAATACATTATGAACCAATAATAGTTGGIGAATTCAAATAICTCCATTAAATTTTTTG 2100

FIG. 10D



2101 AAATCTACAAATTATTAAATATTAGTCAATAACAATGCATAGAAAGTTCCAAAAAAATT 2160
2161 TTGTTAACAGAAACTTCCAAATTTTTTTTTTATGGAACAAGAAATAACAGATAGAAAA 2220
2221 CTATTTTGTGTGGAATGGAAGTAGTAATATACATTAAAGCAAATTTTAAAAAATTATATA 2280
2281 AGCCTATACGGGCTCAAAGTATGTTATCTAGTAGGTGTAATTAATAATGCATGGTGGAT 2340
2341 TCAGAAATTGGGACAACAATGAAAACGGAATTTAAAAATATTAACTTTAAAAATAAATAAAT 2400
2401 TTGAGTAAATGTGTTTTCTGACTATTGAGGGGCAAAAAAGACAATGCCAAAAGTCTAC 2460
2461 GGGTTTGACTGTCCAGTTCGGTAATAATCTAATACTCTGTCTTTGACCGCAGCTCGTG 2520
2521 TAGGGGTCTTCTGACATTTTCACTGTCTTCTACCCCTACTCGTGAGCCCCACCCCTTTTCCCA 2580
2581 TATCCTAAGGGTAATTTTGGAATCCCAATTTAAACCGATTGAGACCGTACCGGACTTCC 2640

FIG. 10E



2641 TGGGATTCTGCTGGAGCATTTATCAAAAATTATTAGCAGCAATGGGTTTATTAAATTTAAA 2700
2701 AACTCACAACTTGATCAGATAAAATTTTCATAAACACTTTTACGATGGATTCGTACGATCT 2760
2761 ATCTAATGACTTTTTTTTTCTACCACGGTGGATGAAAGTTATAGTACTATTAGCCAGAG 2820
2821 ACAATTGATTATAGATAATCCATTATCCATGATATTTAIGATATAAATAGCTGTTAAA 2880
2881 CTATTTTCAGCATCGCAGCTTTCIGCAACTTTTGTTTTTAATTTAAGAGTTTAAATAATAA 2940
2941 AAGTATTAAGGAGCATAACGAGGCAACAAAAGTAATGAACACGGAGAAACAAAAGCCA 3000
3001 TGAAGCTCATTGGTTAGTTAAGCTTAATAAGAAGATTTTATTAAATTTTAAATGACCATG 3060
3061 ATAACAATTATATTTCTGACTTCTTTAAACCCCTCTTACAACAGAAAGCTCCCTTTT 3120
3121 TCAGTAGAAGTCGATTCCCAATCTTAAGACAAAGCCATTAGAAAGAGAAAGTGAGTGA 3180

FIG. 10F



3721 GCACGCTTAGAACTATCTACTTAAAGAATGTTGAGAGACATACTTTGTTGTTCTTGAT 3780
3781 TCITGGCAACAAAACCTCTATTTTCTCTGCTCAAGACGAGGCCAAAAGGTTTTTAATTTT 3840
3841 ATCTTTTATTTTGCTAAATTTTTTTTGTTTATGAATCTTTAGAGTTTCTAACTTTTTTTTTT 3900
3901 TTTAATTGAACAGTTTACGTTTAACTAATGGCGAAGCATATAATGAGTATGGAATCCTGG 3960
3961 AGAAGAAGAAACAGAGCAATTAAAGAAAGAGTAIGTAACTTTTCATGAAGGAGTTGTCTC 4020
4021 TGCTCCTCTAAATCTACCAGGAAGTCTTATCATAAAGCTCTTCAGGTACATTTAATTTT 4080
4081 TTTTGCTGTAAAGTCACAACTCTCATTATAGGTTTTTAAATTTTATGTTAAAT 4140
4141 AAAATATCTAAAATGGTTGTGTAGTCACGAGCAACGATATTGAAGTTTCATTGAGAGGAAA 4200
4201 ATGGAAGAGAGAAAATTGGATATCAAGGAAGAAGATCAAGAAGAAGAAGTGAACAACA 4260

FIG. 10H



4261 GAGGATGAAGCAGAGATGAGTAAGAGTGATCATGTTAGGAAACAAGAACAGACGATGAT 4320
4321 CTTTTGGGATGGGTTTTGAAACATTTCGAATTTATCGACGGAGCAAATTCICGATCICATT 4380
4381 CTTAGTTTGTTATTTGCCGGACATGAGACTTCCTTCTGTAGCCATTGCTCTCGCTATCTTC 4440
4441 TTCTTGCAAGCTTGCCCTAAAGCCGTGAAGAGCTTAGGGTAAGATAATTATAACAGCAC 4500
4501 AAGTTAATTACTACCAAATGTTACGTATTATATAAGTTATTATAGAATTATCTATTAG 4560
4561 AATATACGATGAAAAAGTAIGTATATTTAATTGTCACIAATTTTATGTTTATTGATTIA 4620
4621 TACTTTTGAAGGAAGAGCATCTTGAGATCgCGAGGGCCAAGAAGGAAGGAGAGTCAG 4680
4681 AATTAAATTGGGATGATTACAAGAAAATGGACTTTACTCAATGTGTGTAIGTTACTIATCATT 4740
4741 CTCATTATTCTATGTTTCATATGATTATGATGAACCAAAATTATTGATTTTTTTTTT 4800

FIG. 10I



4801 TTGGTGTGTGAAGTTATAAATGAAACTCTTCGATTCGGAAATGTAGTGGTTTGG 4860
[REDACTED]
4861 CATCGCAAGCACTCAAAGATGTTTCGGTACAAAGGTAAACTTTACGTACAAAAATTTTAA 4920
[REDACTED]
4921 AATAATGAAATCCGGAATATTGAAATCTTATTGGATGAAAAATATTAAAAATAATTACAT 4980
[REDACTED]
4981 TTCTTAATGTTGGAAAAAGGATACGATATCCCTAGTGGTGGAAAGTGTACCGGTGAT 5040
[REDACTED]
5041 CTCAGCCGTACATTTGGATAATTCTCGTTATGACCAACCTAATCTCTTTAATCCTTGGAG 5100
[REDACTED]
5101 ATGGCAACAGGTAAATAAAAAGTTTCTCTCGTTAACTATCGAAAATTAGTGATAGTTTT 5160
[REDACTED]
5161 TTCATCTATTGCAATGATAGATACGTCCTACGTGATTTACCTATCTATAGATACTATACG 5220
[REDACTED]
5221 AGAACTATTAATCTGGCAAAAACTTTTTATTATTATTATCTTTCAAGTTAGATCTTAACA 5280
[REDACTED]
5281 CGTCATGGATCATTCACATGAAAGCATATAAATTAAAAATAAGAGAGAGAAAGAGAC 5340
[REDACTED]

FIG. 10J



5341 GTGTGGGTGAAGTGACGTGAAGACAATAATTAGTAGGATGGTATGCTTTAATGACG 5400
5401 TAGGAGCTGCCTAAATATTCTTATAATCGTGACCGTTGATTTATTATTAGTCACGGCTTT 5460
5461 GATACAAATTTAAGATTTGACGGACGATGGTACCACGGCTTTGACGGATCTCACACGCCCG 5520
5521 ATGACTTGTACGTGGTTAGATTCIGCCACGTTGACTGGTTTAACTTAGATTTATAA 5580
5581 CTCTATTAATTATAACAACATCAAAATCGGGAATTAGAGAAATATACTATATAGTATTA 5640
5641 TTATGATTATTATGAGATAATACTTTATGAAATAAGATAATAATGTTAGTCATGATGTTA 5700
5701 TAGTGAGTGGGAAGGTAAGAGGTGGTGAGAGATGATTAAATGACCCACGTTGGTGGTG 5760
5761 CCAACAAGCACGTGTTCTTCTTCCTTTTTTCTTCCCAACTTCTTTTTTTGGGGGTTTATT 5820
5821 GTGATTATAAAATCGGTTGTGCTTTTTTTTGTGACGAGCAGCAAAACAACGGAGCGT 5880
[REDACTED] exon 8

FIG. 10K



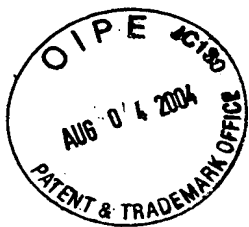
5881 CATCGTCAGGAAGTGGTAGTTTTTCGACGTGGGGAACAACACTACATGCCGTTTGGAGGAG 5940
5941 GGCCAAGGCTATGTGCTGTTAGAGCTAGCCAAGTTAGAAATGGCAGTGTTTATTCATC 6000
6001 ATCTAGTTCCTTAAATTCATTGGGAATTAGCAGAAGATGATCAACCATTGCTTTTCCTT 6060
6061 TTGTTGATTTTCCTAACGGTTTGCCTATTAGGGTTTCTCGTATTCGTAAAAAATAAAAA 6120
6121 AGATGAAAGTATTTTATCT 6180
6181 CCAATGATATATAAAAAATTTGGATAAATAATATTATTGGATATTCGTTTTTTAGTTCGGG 6240
6241 TTTGAGAAAAGGGTTTCGACTTTTCGAAAGTGGACGATGATATAGATTGGGAGCTAGGTT 6300
6301 GAGTCITTTGGACATTGTGATTGGATGTTGTTGATTATTAGTGTGACACTATTAAACCTT 6360
6361 AAATGGGCTTCTATAAGGCCCAATTATATTACGATTATAACAAGTGACAACCTTTTACT 6420

FIG. 10L



6421 TCGTTTTIGATCCGAAGCAATAACAAATTGTCAAATACCAACACACAAGAAATTATGTAAAC 6480
6481 ACTCGTGTGTCTAGTGGGAAATCATTTGGGCTGGAGACTGAACATCAGAACACAAGAAA 6540
6541 CCTGTCAATTATGGATACACCTCCTATGACGGTTTCCAACTTTAICTTTGATTCTTATCG 6600
6601 TGTTACATTGACACAAAGAGTTAGGTGTCAAAGGACTAAATGAATAACAATAGCTCTCA 6660
6661 GGATAAGAAGGTTCATAAAATGGTTCTTTATTTTGAGAAGAAAGAGAGAGGAGCCTTTTA 6720
6721 CTGTTTCTGGGTCCCTATTCCCTTAAATGAGAGGGTTTCGTTTTTACTTCTTCTAICTCA 6780
6781 TCATCTTTAGGATCCCTCTTCTAGACGAGTAAAGTAATCCTCGTTACCAAGCAATGGTCTC 6840
6841 ATCTTTTGAAGACAGGTCCTTTTCCAAAGTCCTAGTTCAGGCCCAAAGCTT 6888

FIG. 10M



1 MFETEHTLL PLLLLPSLLS LLLFLILLKR RNRKTRFNLP PGKSGWPF LG ETIGYLPYT
61 ATTLGDFMQQ HVSKYGKIYR SNLFGEPTIV SADAGLNRFI LQNEGR LFEC SYPRSIGGIL
121 GKWSMLVLVG DMHRDMRSIS LNFLSHARLR TILLK DVERH TLFVLD SWQQ NSIFS AQDEA
181 KKFTFNLMAK HIMSMDPGEE ETEQLKKEYV TFMKGVSAP LNLPGTAYHK ALQSRATILK
241 FIERKMEERK LDIKEEDQEE EEVKTEDEAE MSKSDHVRKQ RTDDDLLGWV LKHSNLSTEQ
301 ILDLILSLF AGHETSSVAI ALAIFFLQAC PKAVEELREE HLEIARAKKE LGESELNWDD
361 YKKMDFTQCV INETLRLGNV VRFLHRKALK DRYKGYDIP SGWKVLPVIS AVHLDNSRYD
421 QPNLFNPWRW QQQNNGASSS GSGSFSTWGN NYMPFGGPR LCAGSELAKL EMAVFIHHLV
481 LKFNWELAE DQPFAPFVD FPNGLP IRVS RIL

FIG. 11

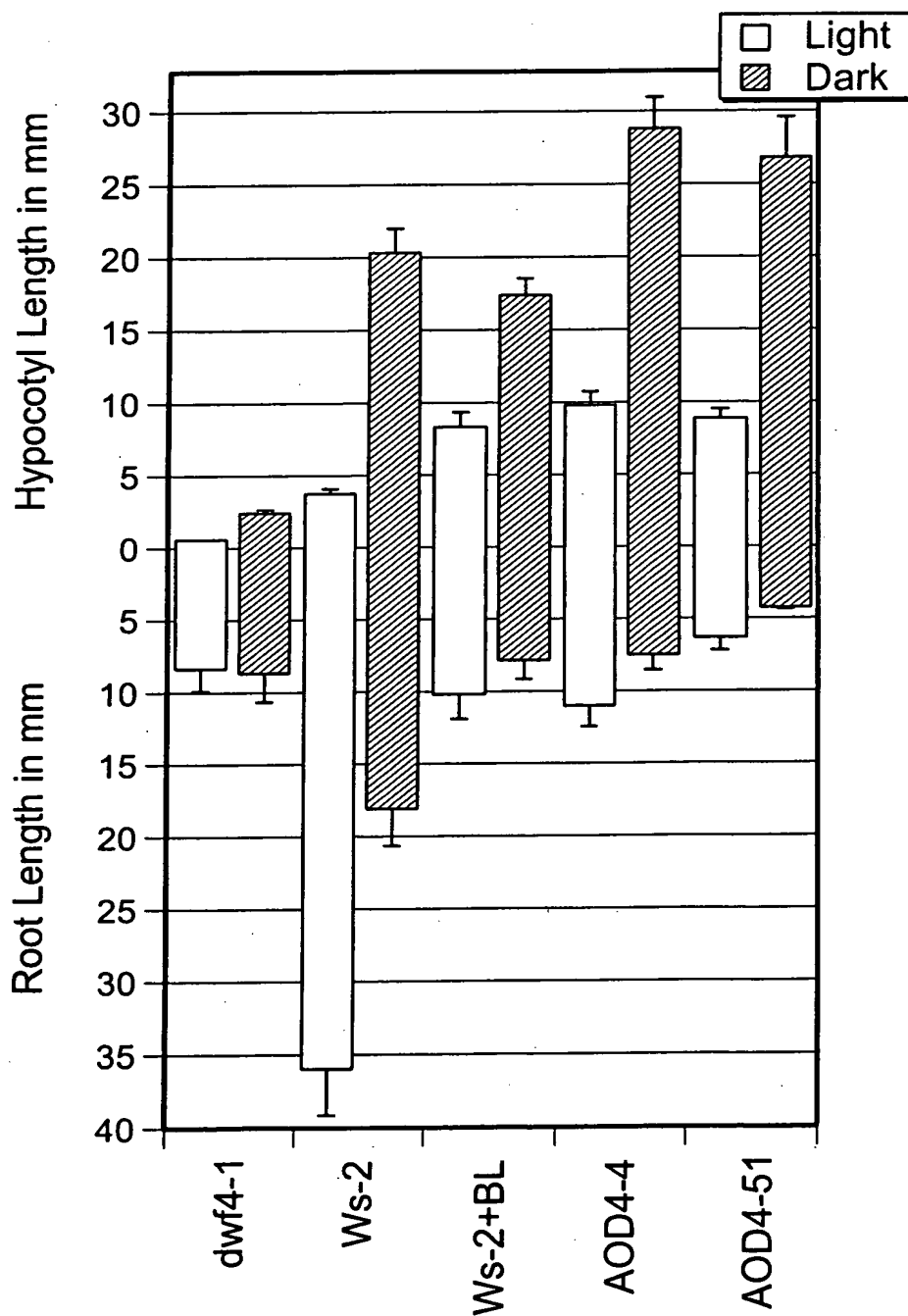


FIG. 12